

USSN 09/551,977

II. REMARKS

Claims 1-37 were pending. Pursuant to a Restriction Requirement, claims 1-16, 18, and 24-37 were withdrawn from consideration.

Applicants gratefully acknowledge Examiner Li's telephone call to discuss the case. During this call, it was agreed that the foregoing amendments to the claims and specification would place the case in condition for allowance.

In particular, the specification has been amended to insert sequence identification numbers were appropriate, particularly on pages 34-35 and 37. Claims 1-16, 18 and 24-37 have been canceled by amendment herein and the dependencies of claims 19, 21, 22 and 23 have been corrected. Applicants reserve the right to file a continuation or divisional application directed to the subject matter of the canceled claims at any time during the pendency of this application. In claim 17, Applicants have replaced the language "comprising one or more" with the equivalent term "comprising an," as described throughout the application as filed, for example, on page 5, lines 4-9 and in the original claims. (See, *PPG Indus v. Guardian Indus Corp.* 48 USPQ2d 1351, 1353 (Fed. Cir. 1998); *Elkay Mfg. Co. v. Ebcø Mfg. Co.*, 52 USPQ2d 1109, 1112, 1353 (Fed. Cir. 1999); *AbTox, Inc. v. Exitron Corp.*, 43 USPQ2d 1545, 1548 (Fed. Cir. 1997); Robert Faber ed., *Landis on Claim Drafting* 531 (3 ed. 1990) establishing that the term "an" in patent parlance carries the meaning of "one or more" in open-ended claims containing the transitional phrase "comprising").

Since the pending claims define an invention that is novel and fully enabled and defined by the specification, Applicant requests that the application proceed to allowance.

Please direct all further communications regarding this application to:

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USSN 09/551,977

Version Showing Changes MadeIn the specification:

The text on starting on line 3 of page 34 and ending on line 7 of page 35 has been amended as follows:

-- Synthesis of cDNA was accomplished by PCR amplification, using the primer sets shown below (Sindbis nucleotide numbering indicated for each primer):

1	CCACAAGCTTGATCTAATGTACCAGCCTGATGC	11472-11450	(SEQ ID NO:3)
1.1	CCACGAATTCAGCCAGATGAGTGAGGC	10364-10381	(SEQ ID NO:4)
2	CCACAAGCTTCAATTCGACGTACGCCTCAC	10394-10375	(SEQ ID NO:5)
2.1	CCACGAATTCATATGGGGAAATCATGAGCC	9614-9634	(SEQ ID NO:6)
3	CCACAAGCTTCATAGACCCTCACTGGCTC	9648-9630	(SEQ ID NO:7)
3.1	CCACGAATTCAGATTAGCACCTCAGGACC	8878-8899	(SEQ ID NO:8)
4	CCACAAGCTTCTACACGGTCCTGAGGTGC	8908-8887	(SEQ ID NO:9)
4.1	CCACGAATTCGTCCGATCATGGATAACTCC	8294-8315	(SEQ ID NO:10)
5	CCACAAGCTTGCGCCACCGAGGAC	8347-8334	(SEQ ID NO:11)
5.1	CCACGAATTCAGTCCATGTGGAGGCC	7797-7814	(SEQ ID NO:12)
6	CCACCTCGAGTTTACCCAACTTAAACAGCC	7368-7348	(SEQ ID NO:13)
6.1	CCACGAGCTCGCGACATTCAATGTGCAATGC	6426-64467	(SEQ ID NO:14)
7	CCACCTCGAGGAACTCCTCCCAATACTCGTC	6488-6468	(SEQ ID NO:15)
7.1	CCACGAGCTCGACCTTGGAGCGCAATGTCC	5843-5862	(SEQ ID NO:16)
8	CCACCTCGAGTTTCGACGTGTCGAGCACC	5900-5882	(SEQ ID NO:17)
8.1	CCACGAGCTCGACCATGGAAGCAATCCGC	4814-4832	(SEQ ID NO:18)
9	CCACCTCGAGACGACGGGTTATGGTCGAC	4864-4845	(SEQ ID NO:19)
9.1	CCACGAGCTCCACGGAGACAGGCACCGC	4246-4264	(SEQ ID NO:20)
10	CCACCTCGAGGATCACTTTCTTTCTAGGCAC	4299-4277	(SEQ ID NO:21)
10.1	CCACGAGCTCGAACTCTCCCGTAGATTTC	3407-3427	(SEQ ID NO:22)

USSN 09/551,977

11 CCACCTCGAGATCAAGTTGTGTGCCCTTCC 3464-3445 (SEQ ID NO:23)
11.1 CCACGAGCTCCCAGGGGATATCATCCTGAC 2742-2761 (SEQ ID NO:24)

12 CCACCTCGAGGCTGTCATTACTTCATGTCCG 2825-2804 (SEQ ID NO:25)
12.1 CCACGAGCTCGAACCGCAAACCTATACCACATTGC 1976-1999 (SEQ ID NO:26)

13 CCACCTCGAGCTTGTACTGCTCCTCTTCTG 2042-2023 (SEQ ID NO:27)
13.1 CCACGAGCTCGGAGAACGGGTATCGTTCC 1029-1047 (SEQ ID NO:28)

14 CCACCTCGAGCCGGGATGTACGTGCAC 1069-1052 (SEQ ID NO:29)
14.1 CCACGAGCTCATTGACGGCGTAGTACACAC 1-20 (SEQ ID NO:30)--

The text on lines 3 through 13 of page 37 has been amended as follows:

--In addition, cDNA clones representing the subgenomic promoter region and 3'-end nontranslated regions also were generated using the following primer pairs:

YSIN1F

5'-GATTCGGTTACTTCCACAGC (SEQ ID NO:31)

YSIN1R

5'-ACTGACGGCTGTGGTCAGTT (SEQ ID NO:32)

YSIN2F

5'-GATGTACTTCCGAGGAACTG (SEQ ID NO:33)

YSIN2R

5'-CCACAAGCTTGAAATGTTAAAAACAAAATTTTGT (SEQ ID NO:34)--

In the claims:

Claims 1-16, 18 and 24-37 have been canceled, without prejudice or disclaimer.

17. (Thrice Amended) A recombinant alphavirus particle which infects human dendritic cells, said recombinant alphavirus particle comprising an [one or more] amino acid [mutations] mutation at about amino acids 158 through 162 of the E2 glycoprotein as compared to wild-type, with the proviso that said recombinant alphavirus particle is not derived from ATCC # VR-2526.

19. (Amended) The recombinant alphavirus particle of claim 17 [or 18] wherein said alphavirus is a Sindbis virus.

USSN 09/551,977

21. (Amended) The recombinant alphavirus particle according to claim 17 [or 18] wherein said alphavirus is Semliki Forest virus.

22. (Amended) The recombinant alphavirus particle according to claim 17 [or 18] wherein said alphavirus is Ross River virus.

23. (Amended) The recombinant alphavirus particle according to claim 17 [or 18] wherein said alphavirus is Venezuelan equine encephalitis virus.

USSN 09/551,977

Currently Pending Claims

1 to 16. Canceled.

17. (Thrice Amended) A recombinant alphavirus particle which infects human dendritic cells, said recombinant alphavirus particle comprising an amino acid mutation at about amino acids 158 through 162 of the E2 glycoprotein as compared to wild-type, with the proviso that said recombinant alphavirus particle is not derived from ATCC # VR-2526.

18. Canceled.

19. (Amended) The recombinant alphavirus particle of claim 17 wherein said alphavirus is a Sindbis virus.

20. The recombinant alphavirus particle according to claim 19 wherein said alphavirus has an amino acid substitution at E2 residue 160, as compared to wild-type Sindbis virus.

21. The recombinant alphavirus particle according to claim 17 wherein said alphavirus is Semliki Forest virus.

22. The recombinant alphavirus particle according to claim 17 wherein said alphavirus is Ross River virus.

23. The recombinant alphavirus particle according to claim 17 wherein said alphavirus is Venezuelan equine encephalitis virus.

24 to 37. Canceled.